

*On page 4, line 25, please replace with:*

--- Target No. 4: Residues 70-77 PDGQSTAK (SEQ. ID. NO. 4)---

*On page 12, lines 29-32, please replace with:*

---The VH and VL genes are linked by an artificial linker (GGGGS)<sub>3</sub> (SEQ.ID.NO. 5) to form scFv fragment [VH-(GGGGS)<sub>3</sub>VL] as in Batra, et al., 1990, and the scFv fragment is subcloned into a pBlueScript vector (Stratagene, CA).---

*On page 13, lines 11-13, please replace with:*

---scFv fragments are linked with the pentamerization domain of COMP via a hinge region (SEQ.ID.NO. 6) to give rise to the following fragment: scFv-hinge [(PQ)<sub>2</sub> PK(PQ)<sub>4</sub> PKPQPK(PE)<sub>2</sub>]- Pentamerization domain (COMP aa 28-72).---

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*On page 18, lines 1-3, please replace with:*

---Sequence carried by the selected phage are then determined using the Sequence kit (United States Biochemical) with the primer 5'-CCCTCATAGTTAAGCGTAACG-3' (SEQ. ID. NO. 15) (Koivunen, E., et al. (1993) J. Bio. Chem 268:20205-20210).---

*On page 18, line 18, please replace with:*

---Target No. 1: VCRHR (SEQ. ID. NO. 7) & GHRCL (SEQ. ID. NO. 8)---

*On page 18, line 19, please replace with:*

---Target No. 2: LGLVTG (SEQ. ID. NO. 9) & RTLVGf (SEQ. ID. NO. 10)---